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A-584.ST25.txt  
SEQUENCE LISTING

<110> TREUHEIT, MICHAEL J.  
O'CONNER, SHEILA R.  
KOSKY, ANDREW A.

<120> PROCESS FOR CORRECTION OF A DISULFIDE MISFOLD IN Fc MOLECULES

<130> A-584

<140> US 09/709,704  
<141> 2000-11-09

<150> US 60/165,188  
<151> 1999-11-12

<160> 6

<170> PatentIn version 3.1

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ggg gga ccg tca gtc ttc ctc ttc ccc cca aaa ccc aag gac acc ctc 96  
Gly Gly Pro Ser Val Phe Leu Phe 25 Pro Lys Pro Lys Asp Thr Leu 30  
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atg atc tcc cgg acc cct gag gtc aca tgc gtg gtg gtg gac gtg agc 144  
Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser 45  
35 40  
cac gaa gac cct gag gtc aag ttc aac tgg tac gtg gac ggc gtg gag 192  
His Glu Asp Pro Glu Val 55 Phe Asn Trp Tyr Val Asp Gly Val Glu 60  
50 55  
gtg cat aat gcc aag aca aag ccg cgg gag gag cag tac aac agc acg 240  
Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr 80  
65 70  
tac cgt gtg gtc agc gtc ctc acc gtc ctg cac cag gac tgg ctg aat 288  
Tyr Arg Val Val Ser Val Leu Thr Val 90 Leu His Gln Asp Trp Leu Asn 95  
85  
ggc aag gag tac aag tgc aag gtc tcc aac aaa gcc ctc cca gcc ccc 336  
Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro 110  
100 105  
atc gag aaa acc atc tcc aaa gcc aaa ggg cag ccc cga gaa cca cag 384  
Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln 125  
115 120  
gtg tac acc ctg ccc cca tcc cgg gat gag ctg acc aag aac cag gtc 432  
Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln Val 140  
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A-584.ST25.txt

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Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val      160
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gag tgg gag agc aat ggg cag ccg gag aac aac tac aag acc acg cct      528
Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro      175
                               165                               170

ccc gtg ctg gac tcc gac ggc tcc ttc ttc ctc tac agc aag ctc acc      576
Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr      180
                               185                               190

gtg gac aag agc agg tgg cag cag ggg aac gtc ttc tca tgc tcc gtg      624
Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val      195
                               200                               205

atg cat gag gct ctg cac aac cac tac acg cag aag agc ctc tcc ctg      672
Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu      210
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His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu
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Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr
65                               70                               75                               80

Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn
85                               90                               95

Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro
100                               105                               110

Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln
115                               120                               125

Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln Val
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A-584.ST25.txt

Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val  
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Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro  
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Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr  
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Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val  
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